

Kindly amend the application without prejudice, without admission, without surrender of subject matter, and without any intention of creating any estoppel as to equivalents as follows:

**IN THE SPECIFICATION:**

Please amend the specification without prejudice, without admission, without surrender of subject matter, and without any intention of creating any estoppel as to equivalents to read as follows:

At Page 15, line 1 please replace the following paragraph as shown on the attached "Version With Markings to" to read as follows:

ST11: AGCCAACATTGCTAAATTGGCGCA (SEQ ID NO:11) (see claim 3, WO 95/00664)

ST15: GGTAGAAATTCCCAGCGGGTACTG (SEQ ID NO:12) (see claim 3, WO 95/00664)

DI Since, however, no amplification, or only insufficient amplification, occurred with that primer pair with a number of strains of subspecies IIIa, IV, V and VI, in those cases the following primers were used for the PCR and sequencing:

ST11: AGCCAACCATTGCTAAATTGGCGCA (see claim 3, WO 95/00664)

ST14: TTTGCGACTATCAGGTTACCGTGG (SEQ ID NO:13) (see claim 3, WO 95/00664).

At Page 19, line 9, please replace the following paragraph as shown on the attached "Version With Markings to" to read as follows:

D2 After the end of the PCR reaction, the amplification products were separated by means of agarose gel electrophoresis and visualised by staining with ethidium bromide. The expected product of 167 bp length (primer combination 1) or of 161 bp length (primer combination 2) was observed in all cases in which DNA of strains of the *Salmonella* genus was present (compare Table 1a), but not in the presence of DNA of other tested bacteria (compare Table 1b). After the end of the run, the DNA contained in the gels was transferred by standard methods to nylon filters and hybridised with the oligonucleotide ST14 (TTTGCGACTATCAGGTTACCGTGG (SEQ ID NO:13) (see claim 3, WO